	CRF Errors Corrected by the STIC Systems Branch CRF Processing Dat: 3/19/
91 IV	Changed a file from non-ASCII to ASCII ENTED Self-reduced by: 40 (STIE
	Changed the margins in cases where the sequence text was "wrapped down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited a format enter in an outronk approach of the account, specifically.
	Edited the Current Application Data section with the actual current number. The number inputted by th applicant was \(\) the prior application data; or \(\) other \(\)
	Added the mandatory heading and subheadings for *Current Application Data*.
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: Gron-ASCII "garbage" at the beginning/end of files; Gecretary initials/filename at end o page numbers throughout text; dother invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
_	Corrected an error in the Number of Sequences field, specifically:
_	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted <i>ending</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (en tue to a Patentin bug). Sequences corrected:
	Other:
-	

*Examin r: Th above corrections must b communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



OIPE

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/974,026 TIME: 12:42:21

DATE: 03/19/2002

Input Set : A:\PTO.AMC.txt Output Set: N:\CRF3\03192002\I974026.raw

3 <110> APPLICANT: Tamburini, Paul P Davis, Gary 5 Delaria, Katherine A Christopher, Marlor W Daniel, Muller K 9 <120> TITLE OF INVENTION: Human Bikunin 11 <130> FILE REFERENCE: 96-223-ZZ 13 <140> CURRENT APPLICATION NUMBER: US 09/974,026 14 <141> CURRENT FILING DATE: 2001-10-10 16 <150> PRIOR APPLICATION NUMBER: US 09/144,428 17 <151> PRIOR FILING DATE: 1998-08-31 19 <150> PRIOR APPLICATION NUMBER: PCT/US97/03894 20 <151> PRIOR FILING DATE: 1997-03-10 22 <150> PRIOR APPLICATION NUMBER: US 08/725,251 23 <151> PRIOR FILING DATE: 1996-10-04 25 <150> PRIOR APPLICATION NUMBER: US 60/019,793 26 <151> PRIOR FILING DATE: 1996-06-14 28 <150> PRIOR APPLICATION NUMBER: US 60/013,106 29 <151> PRIOR FILING DATE: 1996-03-11 31 <160> NUMBER OF SEQ ID NOS: 105 33 <170> SOFTWARE: PatentIn version 3.1 35 <210> SEO ID NO: 1 36 <211> LENGTH: 179 37 <212> TYPE: PRT 38 <213> ORGANISM: Homo sapiens 40 <400> SEOUENCE: 1 42 Ala Asp Arg Glu Arg Ser Ile His Asp Phe Cys Leu Val Ser Lys Val 43 1 5 10 46 Val Gly Arg Cys Arg Ala Ser Met Pro Arg Trp Trp Tyr Asn Val Thr 47 20 25 50 Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly Gly Cys Asp Gly Asn Ser 51 35 40 54 Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu Lys Lys Cys Ala Thr Val 58 Thr Glu Asn Ala Thr Gly Asp Leu Ala Thr Ser Arg Asn Ala Ala Asp 70 75 62 Ser Ser Val Pro Ser Ala Pro Arg Arg Gln Asp Ser Glu Asp His Ser 66 Ser Asp Met Phe Asn Tyr Glu Glu Tyr Cys Thr Ala Asn Ala Val Thr 67 100 105

70 Gly Pro Cys Arg Ala Ser Phe Pro Arg Trp Tyr Phe Asp Val Glu Arg

74 Asn Ser Cys Asn Asn Phe Ile Tyr Gly Gly Cys Arg Gly Asn Lys Asn

71 115 120

DATE: 03/19/2002

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/974.026 TIME: 12:42:21

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78 Ser Tyr Arg Ser Glu Glu Ala Cys Met Leu Arg Cys Phe Arg Gln Gln
79 145 150
                            155
82 Glu Asn Pro Pro Leu Pro Leu Gly Ser Lys Val Val Val Leu Ala Gly
             165
                                170
86 Ala Val Ser
90 <210> SEO ID NO: 2
91 <211> LENGTH: 197
92 <212> TYPE: PRT
93 <213> ORGANISM: Homo sapiens
95 <220> FEATURE:
96 <221> NAME/KEY: SIGNAL
97 <222> LOCATION: (1)..(18)
98 <223> OTHER INFORMATION:
101 <400> SEOUENCE: 2
103 Ala Gly Ser Phe Leu Ala Trp Leu Gly Ser Leu Leu Ser Gly Val
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107 Leu Ala Ala Asp Arg Glu Arg Ser Ile His Asp Phe Cys Leu Val Ser
108 20
                               25
111 Lys Val Val Gly Arg Cys Arg Ala Ser Met Pro Arg Trp Trp Tyr Asn
112 35
                           40
115 Val Thr Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly Gly Cys Asp Gly
                       55
119 Asn Ser Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu Lys Lys Cys Ala
                    70
123 Thr Val Thr Glu Asn Ala Thr Gly Asp Leu Ala Thr Ser Arg Asn Ala
                                  90
                85
127 Ala Asp Ser Ser Val Pro Ser Ala Pro Arg Arg Gln Asp Ser Glu Asp
128 100
                               105
131 His Ser Ser Asp Met Phe Asn Tyr Glu Glu Tyr Cys Thr Ala Asn Ala
132 115 120
135 Val Thr Gly Pro Cys Arg Ala Ser Phe Pro Arg Trp Tyr Phe Asp Val
136 130 135
139 Glu Arg Asn Ser Cys Asn Asn Phe Ile Tyr Gly Gly Cys Arg Gly Asn
140 145 150
                                     155
143 Lys Asn Ser Tyr Arg Ser Glu Glu Ala Cys Met Leu Arg Cys Phe Arg
144 165
                                170
147 Gln Gln Glu Asn Pro Pro Leu Pro Leu Gly Ser Lys Val Val Leu
148 180
                              185
151 Ala Gly Ala Val Ser
152 195
155 <210> SEO ID NO: 3
156 <211> LENGTH: 153
157 <212> TYPE: PRT
158 <213> ORGANISM: Homo sapiens
160 <400> SEOUENCE: 3
162 Ile His Asp Phe Cys Leu Val Ser Lys Val Val Gly Arg Cys Arg Ala
163 1 5
                                  10
166 Ser Met Pro Arg Trp Trp Tyr Asn Val Thr Asp Gly Ser Cys Gln Leu
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RAW SEQUENCE LISTING DATE: 03/19/2002 PATENT APPLICATION: US/09/974,026 TIME: 12:42:21

```
25
170 Phe Val Tyr Gly Gly Cys Asp Gly Asn Ser Asn Asn Tyr Leu Thr Lys
171 35
                           40
174 Glu Glu Cys Leu Lys Lys Cys Ala Thr Val Thr Glu Asn Ala Thr Gly
                         55
                                            60
178 Asp Leu Ala Thr Ser Arg Asn Ala Ala Asp Ser Ser Val Pro Ser Ala
                     70
                                        75
182 Pro Arg Arg Gln Asp Ser Glu Asp His Ser Ser Asp Met Phe Asn Tyr
                                    90
186 Glu Glu Tyr Cys Thr Ala Asn Ala Val Thr Gly Pro Cys Arg Ala Ser
            100
                                 105
190 Phe Pro Arg Trp Tyr Phe Asp Val Glu Arg Asn Ser Cys Asn Asn Phe
191 115
                            120
194 Ile Tyr Gly Gly Cys Arg Gly Asn Lys Asn Ser Tyr Arg Ser Glu Glu
195 130
               135
                                           140
198 Ala Cys Met Leu Arg Cys Phe Arg Gln
199 145
202 <210> SEO ID NO: 4
203 <211> LENGTH: 58
204 <212> TYPE: PRT
205 <213> ORGANISM: Homo sapiens
207 <400> SEQUENCE: 4
209 Ile His Asp Phe Cys Leu Val Ser Lys Val Val Gly Arg Cys Arg Ala
210 1 5
213 Ser Met Pro Arg Trp Trp Tyr Asn Val Thr Asp Gly Ser Cys Gln Leu
217 Phe Val Tyr Gly Gly Cys Asp Gly Asn Ser Asn Asn Tyr Leu Thr Lys
                         40
221 Glu Glu Cys Leu Lys Lys Cys Ala Thr Val
222 50
225 <210> SEQ ID NO: 5
226 <211> LENGTH: 51
227 <212> TYPE: PRT
228 <213> ORGANISM: Homo sapiens
230 <400> SEOUENCE: 5
232 Cys Leu Val Ser Lys Val Val Gly Arg Cys Arg Ala Ser Met Pro Arg
233 1
                                    10
236 Trp Trp Tyr Asn Val Thr Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly
237 20
                                25
                                                 30
240 Gly Cys Asp Gly Asn Ser Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu
241 35
                            4.0
244 Lys Lys Cys
245 50
248 <210> SEO ID NO: 6
249 <211> LENGTH: 58
250 <212> TYPE: PRT
251 <213> ORGANISM: Homo sapiens
253 <400> SEOUENCE: 6
255 Tyr Glu Glu Tyr Cys Thr Ala Asn Ala Val Thr Gly Pro Cys Arg Ala
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DATE: 03/19/2002

TIME: 12:42:21

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/974,026

```
259 Ser Phe Pro Arg Trp Tyr Phe Asp Val Glu Arg Asn Ser Cys Asn Asn
260 20
                                 25
263 Phe Ile Tyr Gly Gly Cys Arg Gly Asn Lys Asn Ser Tyr Arg Ser Glu
264 35
                             40
                                           4.5
267 Glu Ala Cys Met Leu Arg Cys Phe Arg Gln
268 50
                          55
271 <210> SEO ID NO: 7
272 <211> LENGTH: 51
273 <212> TYPE: PRT
274 <213> ORGANISM: Homo sapiens
276 <400> SEQUENCE: 7
278 Cys Thr Ala Asn Ala Val Thr Gly Pro Cys Arg Ala Ser Phe Pro Arg
279 1 5
                                  10
282 Trp Tyr Phe Asp Val Glu Arg Asn Ser Cys Asn Asn Phe Ile Tyr Gly
283 20
                                  25
286 Gly Cys Arg Gly Asn Lys Asn Ser Tyr Arg Ser Glu Glu Ala Cys Met
287 35
                           40
290 Leu Arg Cys
291 50
294 <210> SEO ID NO: 8
295 <211> LENGTH: 92
296 <212> TYPE: PRT
297 <213> ORGANISM: Homo sapiens
299 <400> SEQUENCE: 8
301 Ala Asp Arg Glu Arg Ser Ile His Asp Phe Cys Leu Val Ser Lys Val
                                     1.0
305 Val Gly Arg Cys Arg Ala Ser Met Pro Arg Trp Trp Tyr Asn Val Thr
              20
                                  25
309 Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly Gly Cys Asp Gly Asn Ser
310 35
                              40
313 Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu Lys Lys Cys Ala Thr Val
                          55
317 Thr Glu Asn Ala Thr Gly Asp Leu Ala Thr Ser Arg Asn Ala Ala Asp
                    70
321 Ser Ser Val Pro Ser Ala Pro Arg Arg Gln Asp Ser
325 <210> SEO ID NO: 9
326 <211> LENGTH: 708
327 <212> TYPE: DNA
328 <213> ORGANISM: Artificial Sequence
330 <220> FEATURE:
331 <223> OTHER INFORMATION: Consensus DNA sequence of human Bikunin (Fig. 3).
333 <220> FEATURE:
334 <221> NAME/KEY: misc feature
335 <222> LOCATION: (679)..(679)
336 <223> OTHER INFORMATION: "n" is any nucleotide.
339 <220> FEATURE:
340 <221> NAME/KEY: misc_feature
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RAW SEQUENCE LISTING

DATE: 03/19/2002 PATENT APPLICATION: US/09/974.026 TIME: 12:42:21

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341 <222> LOCATION: (707)..(707)
    342 <223> OTHER INFORMATION: "n" is any nucleotide.
    345 <400> SEOUENCE: 9
    346 ggccgggtcg tttctcgcct ggctgggatc gctgctcctc tctggggtcc tggcggccga
                                                                              60
    348 cogagaacgc agcatccacg acttctgcct ggtgtcgaag gtggtgggca gatgccgggc
                                                                             120
    350 ctccatgcct aggtggtggt acaatgtcac tgacggatcc tgccagctgt ttgtgtatgg
                                                                             180
    352 gggctgtgac ggaaacagca ataattacct gaccaaggag gagtgcctca agaaatgtgc
                                                                             240
    354 cactotcaca gagaatgcca cgggtgacct ggccaccagc aggaatgcag cggattcctc
                                                                             300
    356 tgtcccaagt gctcccagaa ggcaggattc tgaagaccac tccagcgata tgttcaacta
                                                                             360
    358 tgaaqaatac tgcaccgcca acgcaqtcac tgggccttgc cgtgcatcct tcccacgctg
                                                                             420
    360 gtactttgac gtggagagga actectgcaa taacttcate tatggagget geeggggcaa
                                                                             480
    362 taaqaacaqc taccqctctq aqqaqqcctq catqctccqc tqcttccqcc aqcaqqaqaa
                                                                             540
                                                                             600
    364 tectecety eccettaget casaggtggt ggttetggee ggggetgttt cgtgatggtg
    366 ttgatcettt teetggggag catecatggt ettactgatt eegggtggca aggaggaace
                                                                             660
w W 368 aggagegtge cetgeggane gtetggaget teggagatga caagggnt
                                                                             708
    371 <210> SEO ID NO: 10
    372 <211> LENGTH: 197
    373 <212> TYPE: PRT
    374 <213> ORGANISM: Artificial Sequence
    376 <220> FEATURE:
    377 <223> OTHER INFORMATION: Amino acids -18 to 179 of translation of consensus sequence
in Fig. 3.
    379 <400> SEQUENCE: 10
    381 Ala Gly Ser Phe Leu Ala Trp Leu Gly Ser Leu Leu Ser Gly Val
                                            1.0
    385 Leu Ala Ala Asp Arg Glu Arg Ser Ile His Asp Phe Cys Leu Val Ser
                   20
                                        25
     389 Lys Val Val Gly Arg Cys Arg Ala Ser Met Pro Arg Trp Trp Tyr Asn
               35
                                    40
     393 Val Thr Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly Gly Cys Asp Gly
                                55
    397 Asn Ser Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu Lys Lys Cys Ala
                            70
                                                75
    401 Thr Val Thr Glu Asn Ala Thr Gly Asp Leu Ala Thr Ser Arg Asn Ala
                        85
                                            90
    405 Ala Asp Ser Ser Val Pro Ser Ala Pro Arg Arg Gln Asp Ser Glu Asp
                   100
                                        105
    409 His Ser Ser Asp Met Phe Asn Tyr Glu Glu Tyr Cys Thr Ala Asn Ala
               115
                                    120
    413 Val Thr Gly Pro Cys Arg Ala Ser Phe Pro Arg Trp Tyr Phe Asp Val
                                135
                                                   140
     417 Glu Arg Asn Ser Cys Asn Asn Phe Ile Tyr Gly Gly Cys Arg Gly Asn
                            150
                                                155
     421 Lys Asn Ser Tyr Arg Ser Glu Glu Ala Cys Met Leu Arg Cys Phe Arg
                                           170
                       165
    425 Gln Gln Glu Asn Pro Pro Leu Pro Leu Gly Ser Lys Val Val Val Leu
                   180
                                        185
    429 Ala Gly Ala Val Ser
               195
    433 <210> SEO ID NO: 11
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/974.026 DATE: 03/19/2002 TIME: 12:42:22

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\03192002\1974026.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220 to <223) fields of each sequence which presents at least one n or Xaa.

```
Seg#:9; N Pos. 679,707
Seq#:11; Xaa Pos. 8,17,19,21,22,23,24,25,26,40,42,45,46,47,52,64,103,112
Seq#:11; Xaa Pos. 114,116,117,118,119,120,121,135,137,140,141,142,147,159
Seg#:12; N Pos. 361,367,384,390
Seg#:14: N Pos. 424,481,509
Seg#:16: N Pos. 3,11,12,17,48,425
Seg#:17; N Pos. 7,403,409
Seg#:48; N Pos. 1358
Seg#:51; N Pos. 46,117,313
Seg#:72: Xaa Pos. 9,11,17,19
Seg#:74; Xaa Pos. 25
Seq#:75; N Pos. 425,482,510
Seq#:76; Xaa Pos. 25
Seg#:77; N Pos. 45,49,118,231,305
Seg#:78: N Pos. 117,123,321
Seg#:79; N Pos. 9,11,222,231,262,267,274
Seg#:80: N Pos. 44.46.76.114.187.268.309.317.332.370
Seg#:81: N Pos. 35,148,235,261,272,293,300,313,320
Seg#:82; N Pos. 56,137,145,159,233
Seq#:83; N Pos. 20,26,95,292,313,314,315
Seq#:84; N Pos. 27,139,223,232,302,310,322,328,357,375,392,398,405,427,437
Seg#:84: N Pos. 449,458,474
Seg#:85; N Pos. 361,367,384,390
Seg#:86; N Pos. 3,11,12,17,48,425
Seq#:87; N Pos. 7,403,409
Seq#:88; N Pos. 48,62,211,232,245,309,318
Seg#:89; N Pos. 424,481,509
Seq#:90; N Pos. 257
Seq#:91; N Pos. 19,147
Seg#:92; N Pos. 33,55,213,228,259,267,324,333,344,387
Seq#:93; N Pos. 306,328,342,365,370,377,382,402
Seq#:94; N Pos. 1,142,339,347
Seq#:95; N Pos. 334,368,376
Seg#:96; N Pos. 108,261
Seg#:97; N Pos. 20,30
Seq#:98; N Pos. 45,102,105,159,174,213,337
Seg#:100; N Pos. 304,309
Seq#:101; N Pos. 24
Seq#:102; N Pos. 61,74,122,184
Seq#:103; N Pos. 7
Seq#:104; N Pos. 32,67,136
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Seq#:105; N Pos. 13,19,107



OTPE

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/974,026

DATE: 03/11/2002 TIME: 15:03:12

Input Set : A:\09-974.026 sequence listing.txt

Output Set: N:\CRF3\03112002\I974026.raw

may est Does Not Comply Corrected Diskette Needed

```
3 <110> APPLICANT: Tamburini, Paul P
        Davis, Gary
5
        Delaria, Katherine A
        Christopher, Marlor W
        Daniel, Muller K
9 <120> TITLE OF INVENTION: Human Bikunin
11 <130> FILE REFERENCE: 96-223-ZZ
13 <140> CURRENT APPLICATION NUMBER: US 09/974,026
14 <141> CURRENT FILING DATE: 2001-10-10
16 <150> PRIOR APPLICATION NUMBER: US 09/144,428
17 <151> PRIOR FILING DATE: 1998-08-31
19 <150> PRIOR APPLICATION NUMBER: PCT/US97/03894
20 <151> PRIOR FILING DATE: 1997-03-10
22 <150> PRIOR APPLICATION NUMBER: US 08/725,251
23 <151> PRIOR FILING DATE: 1996-10-04
25 <150> PRIOR APPLICATION NUMBER: US 60/019.793
26 <151> PRIOR FILING DATE: 1996-06-14
28 <150> PRIOR APPLICATION NUMBER: US 60/013,106
29 <151> PRIOR FILING DATE: 1996-03-11
31 <160> NUMBER OF SEO ID NOS: 105
33 <170> SOFTWARE: PatentIn version 3.1
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ERRORED SEQUENCES

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    3914 <211> LENGTH: 343
    3915 <212> TYPE: DNA
    3916 <213> ORGANISM: Homo sapiens
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    3919 <221> NAME/KEY: misc_feature
    3920 <222> LOCATION: (13)..(13)
    3921 <223> OTHER INFORMATION: "n" is any nucleotide.
    3924 <220> FEATURE:
    3925 <221> NAME/KEY: misc_feature
    3926 <222> LOCATION: (19)..(19)
    3927 <223> OTHER INFORMATION: "n" is any nucleotide.
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    3931 <221> NAME/KEY: misc_feature
    3932 <222> LOCATION: (107)..(107)
    3933 <223> OTHER INFORMATION: "n" is any nucleotide.
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3/11/02

60

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/974,026

DATE: 03/11/2002 TIME: 15:03:12

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Output Set: N:\CRF3\03112002\I974026.raw

						,		
W>	3939	gcatggcctg	cagtctggca	gcagccccga	gttgtttcct	cgctgancga	tttctttcct	120
	3941	ccaggtagag	ttttctttgc	ttatgttgaa	ttccattgcc	tcttttctca	tcacagaagt	180
	3943	gatgttggaa	tcgtttcttt	tgtttgtctg	atttatggtt	tttttaagta	taaacaaaag	240
	3945	ttttttatta	gcattctgaa	agaaggaaag	taaaatgtac	aagtttaata	aaaaggggcc	300
	3947	tyccccttta	gaataaaaaa	aaaaaaaaa	aaaaaaaaa	aaa		343

 $E - \rightarrow 3950 \begin{pmatrix} \hat{1} \\ E - - > 3952 \end{pmatrix} \begin{pmatrix} \hat{1} \\ 1 \end{pmatrix}$

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/974,026

DATE: 03/11/2002 TIME: 15:03:13

Input Set : A:\09-974,026 sequence listing.txt Output Set: N:\CRF3\03112002\I974026.raw

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T::605 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:11
T::609 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:11
L:613 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:617 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:11
L:629 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:11
T.: 633 M: 341 W: (46) "n" or "Xaa" used, for SEO ID#:11
L:637 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:11
L:641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:695 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:12
L:777 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:779 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
T.: 833 M: 341 W: (46) "n" or "Xaa" used, for SEO ID#:16
L:847 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:874 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:886 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:17
L:1731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48
L:1891 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:1893 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:1901 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:2409 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:72
L:2413 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72
L:2469 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74
L:2514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75
L:2516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75
I::2539 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:76
L:2579 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:77
L:2581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77
L:2585 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77
L:2589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77
L:2618 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:2620 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:2626 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:2671 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79
L:2677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79
L:2679 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79
L:2748 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80
L:2750 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80
L:2754 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80
L:2756 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80
L:2758 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80
L:2760 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80
L:2825 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:2829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:2831 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:2833 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:2835 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:2874 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/974,026 TIME: 15:03:13

DATE: 03/11/2002

Input Set : A:\09-974,026 sequence listing.txt Output Set: N:\CRF3\03112002\1974026.raw

L:2878 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 L:2880 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 L:3950 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:343 SEQ:105

M:254 Repeated in SeqNo=105